Biomedical Computing Interest Group



From terminology integration to information integration

An example in the domain of genomics



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Outline

- **♦** Background
 - Terminology integration: The Unified Medical Language System
 - Information integration:
 Genomics as an example
- **♦** Applications
 - GenesTrace
 - BioMeKe



Terminology integration

The Unified Medical Language System

Motivation

- ◆ Started in 1986
- ◆ National Library of Medicine
- ◆ "Long-term R&D project"
- Complementary to IAIMS

(Integrated Academic Information Management Systems)

- «[...] the UMLS project is an effort to overcome two significant barriers to effective retrieval of machine-readable information.
- The first is the variety of ways the same concepts are expressed in different machine-readable sources and by different people.
- The second is the distribution of useful information among many disparate databases and systems.»



Source Vocabularies

(2005AA)

- ◆ 134 source vocabularies
 - 132 contributing concept names
- ◆ ~80 families of vocabularies
 - multiple translations (e.g., MeSH, ICPC, ICD-10)
 - variants (American-English equivalents, Australian extension/adaptation)
 - subsequent editions usually considered distinct families (ICD: 9-10; DSM: IIIR-IV)
- ◆ Broad coverage of biomedicine
- Common presentation



Biomedical terminologies

- ◆ General vocabularies
 - anatomy (UWDA, Neuronames)
 - drugs (RxNorm, First DataBank, Micromedex)
 - medical devices (UMD, SPN)
- Several perspectives
 - clinical terms (SNOMED CT)
 - information sciences (MeSH, CRISP)
 - administrative terminologies (ICD-9-CM, CPT-4)
 - data exchange terminologies (HL7, LOINC)

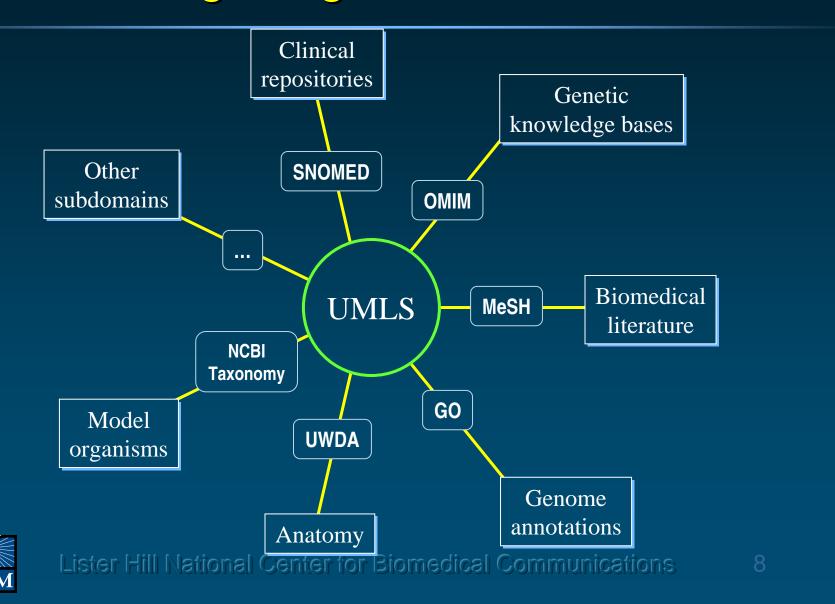


Biomedical terminologies (cont'd)

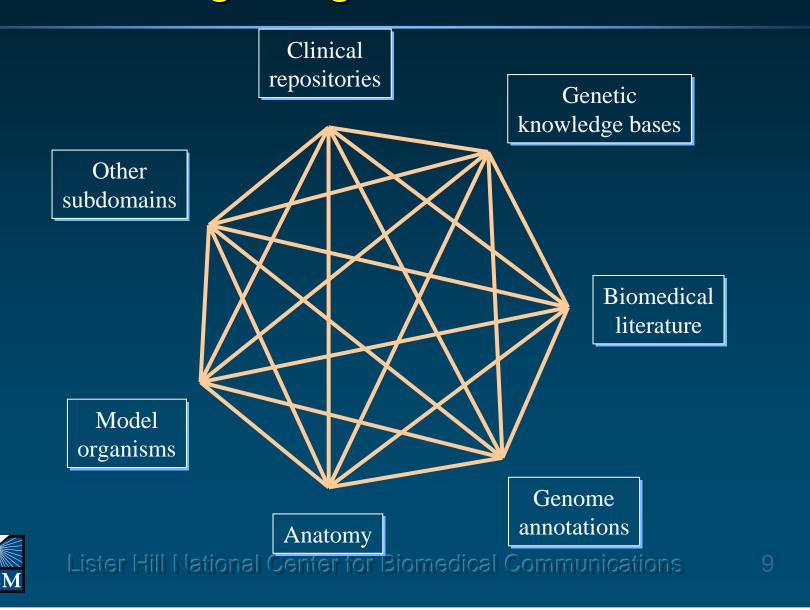
- ◆ Specialized vocabularies
 - nursing (NIC, NOC, NANDA, Omaha, PCDS)
 - dentistry (CDT)
 - psychiatry (DSM, APA)
 - adverse reactions (COSTART, WHO ART)
 - primary care (ICPC)
 - genomics (GO, OMIM, HUGO)
- ◆ Terminology of knowledge bases (AI/Rheum, DXplain, QMR)

The UMLS serves as a vehicle for the regulatory standards (HIPAA, CHI)

Integrating subdomains



Integrating subdomains

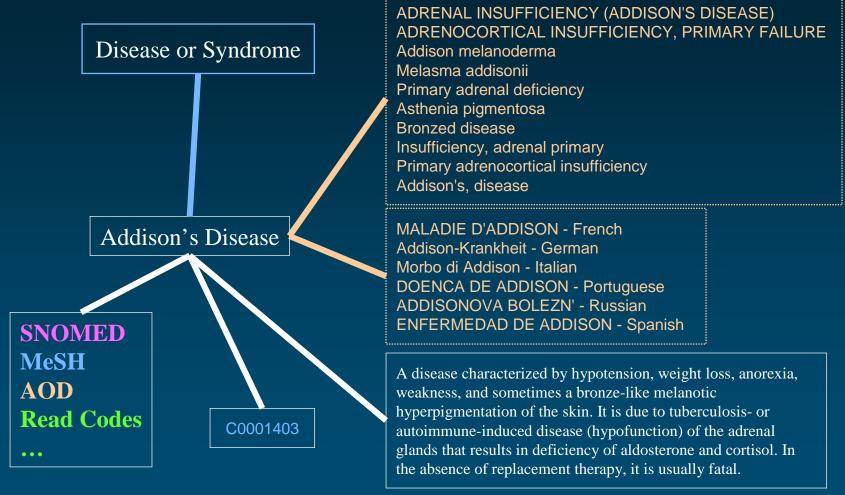


UMLS: 3 components

- ◆ Metathesaurus
 - Concepts
 - Inter-concept relationships
- **♦** Semantic Network
 - Semantic types
 - Semantic network relationships
- **♦** Lexical resources
 - SPECIALIST Lexicon
 - Lexical tools



Addison's Disease: Concept





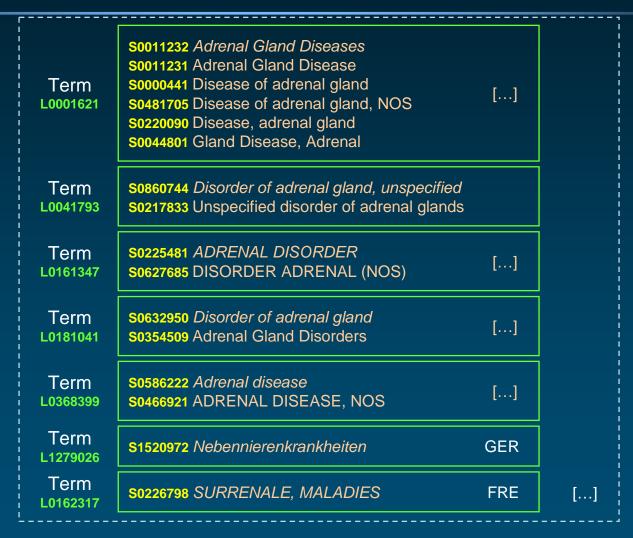
Metathesaurus Concepts (2005AA)

- ◆ Concept (~ 1.2M) CUI
 - Set of synonymous concept names
- ◆ Term (~ 4.2 M) LUI
 - Set of normalized names
- **♦** String (~ 4.7M) **SUI**
 - Distinct concept name
- ◆ Atom (~ 5.5M) AUI
 - Concept name in a given source

```
A0000001 headache
                    (source 1)
A0000002 headache
                    (source 2)
          S0000001
A0000003 Headache (source 1)
A0000004 Headache (source 2)
          S0000002
          L0000001
A0000005 Cephalgia (source 1)
          S000003
          L0000002
          C0000001
```



Cluster of synonymous terms





Concept

C0001621

Metathesaurus Evolution over time

- ◆ Concepts never die (in principle)
 - CUIs are permanent identifiers
- ◆ What happens when they do die (in reality)?
 - Concepts can merge or split
 - Resulting in new concepts and deletions





Metathesaurus Relationships

- ◆ Symbolic relations: ~9 M pairs of concepts
- ◆ Statistical relations : ~7 M pairs of concepts (co-occurring concepts)
- ◆ Mapping relations: 100,000 pairs of concepts

◆ Categorization: Relationships between concepts and semantic types from the Semantic Network



Symbolic relations

- **♦** Relation
 - Pair of "atom" identifiers
 - Type
 - Attribute (if any)
 - List of sources (for type and attribute)
- ◆ Semantics of the relationship: defined by its type [and attribute]

Source transparency: the information is recorded at the "atom" level



Symbolic relationships Type

◆ Hierarchical

Parent / Child PAR / CHD

• Broader / Narrower than RB/RN



◆ Derived from hierarchies

• Siblings (children of parents) SIB



♦ Associative

• Other RO



◆ Various flavors of near-synonymy

• Similar RL

Source asserted synonymy

Possible synonymy RQ

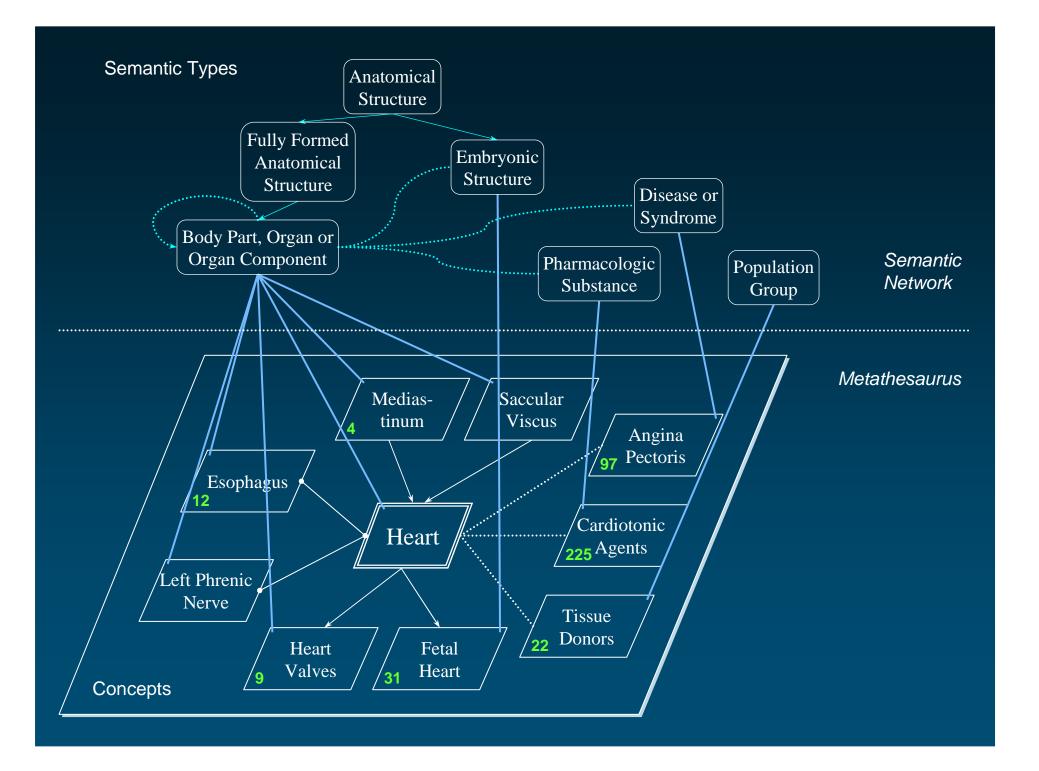




Symbolic relationships Attribute

- **♦** Hierarchical
 - isa (is-a-kind-of)
 - part-of
- **♦** Associative
 - location-of
 - caused-by
 - treats
 - ...
- **◆** Cross-references (mapping)



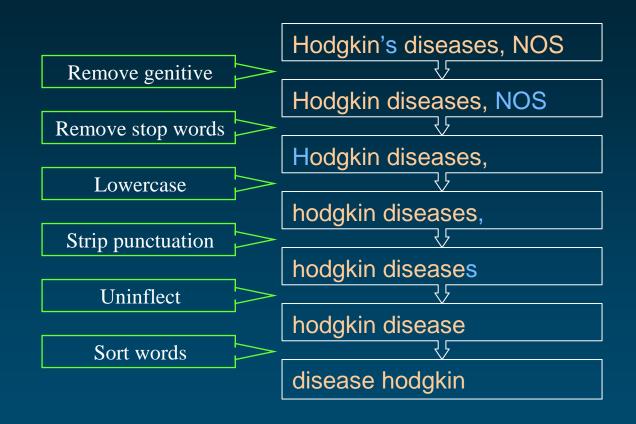


Lexical tools

- ◆ To manage lexical variation in biomedical terminologies
- ◆ Major tools
 - Normalization
 - Indexes
 - Lexical Variant Generation program (lvg)
- ◆ Based on the SPECIALIST Lexicon
- ◆ Used by noun phrase extractors, search engines



Normalization





Normalization: Example

Hodgkin Disease HODGKINS DISEASE Hodgkin's Disease Disease, Hodgkin's Hodgkin's, disease HODGKIN'S DISEASE Hodgkin's disease Hodgkins Disease Hodgkin's disease NOS Hodgkin's disease, NOS Disease, Hodgkins Diseases, Hodgkins **Hodgkins Diseases** Hodgkins disease hodgkin's disease Disease, Hodgkin

normalize disease hodgkin



Information integration

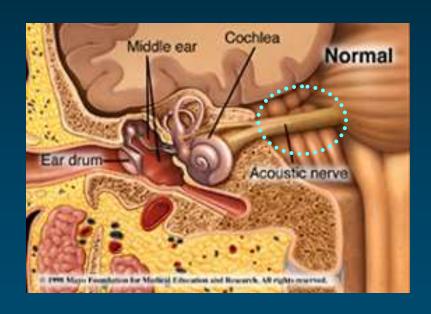
Genomics as an example

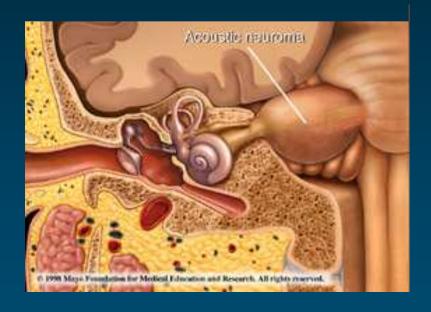
NF2 Gene, protein, and disease

Neurofibromatosis 2 is an autosomal dominant disease characterized by tumors called schwannomas involving the acoustic nerve, as well as other features. The disorder is caused by mutations of the NF2 gene resulting in absence or inactivation of the protein product. The protein product of NF2 is commonly called merlin (but also neurofibromin 2 and schwannomin) and functions as a tumor suppressor.



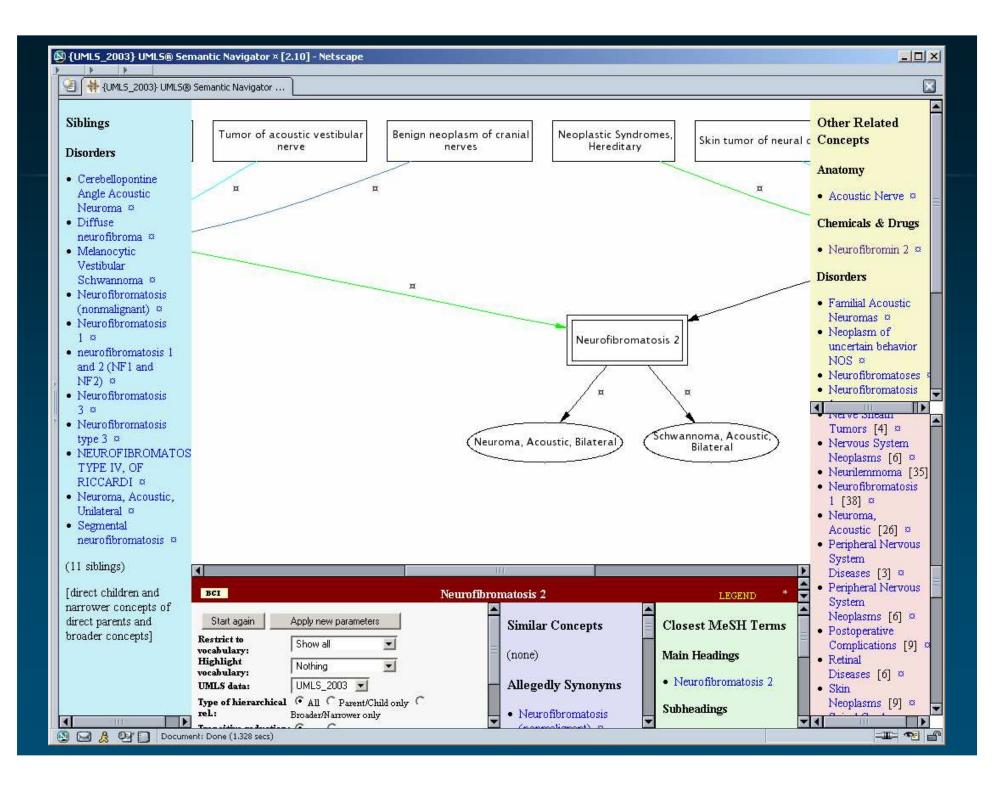
Schwannoma (acoustic neuroma)



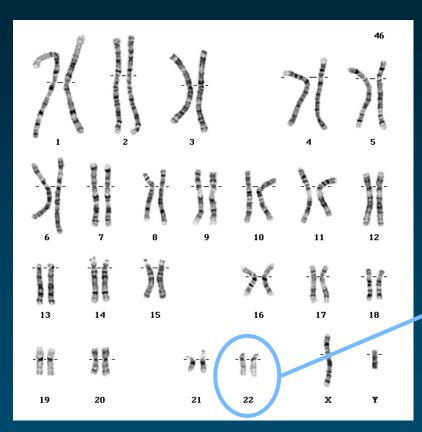


http://www.mayoclinic.com

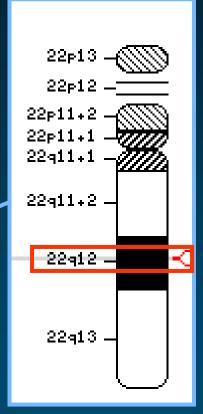




NF2 gene

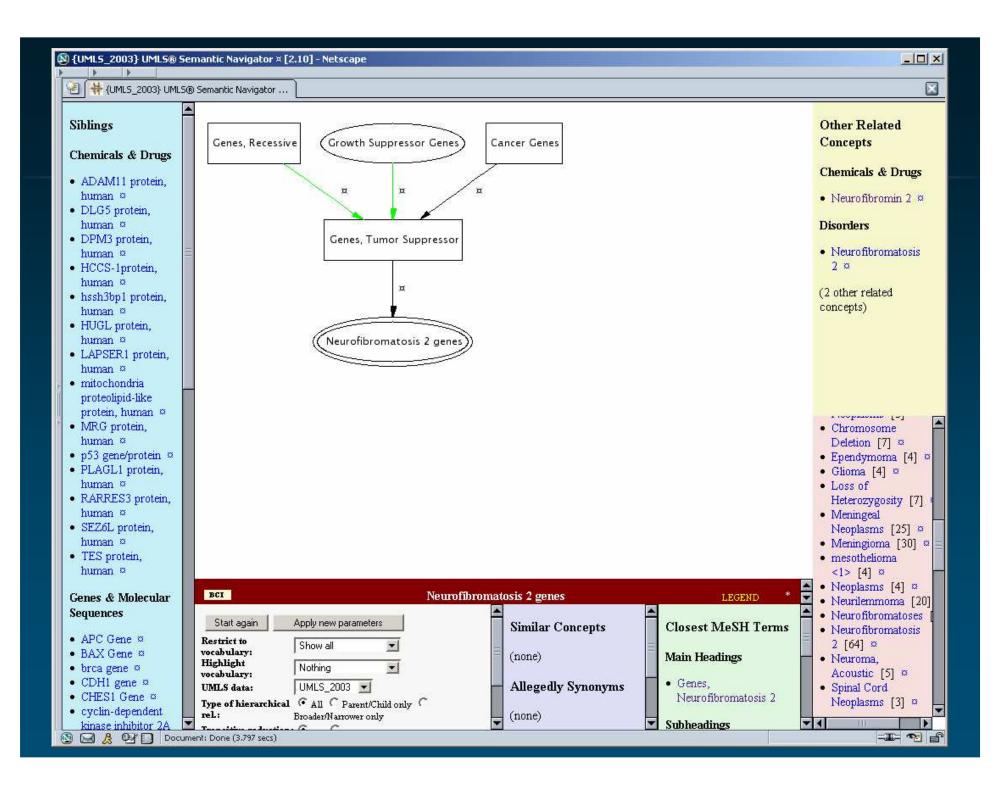


http://staff.washington.edu/timk/cyto/human/



http://www.ncbi.nlm.nih.gov/mapview/

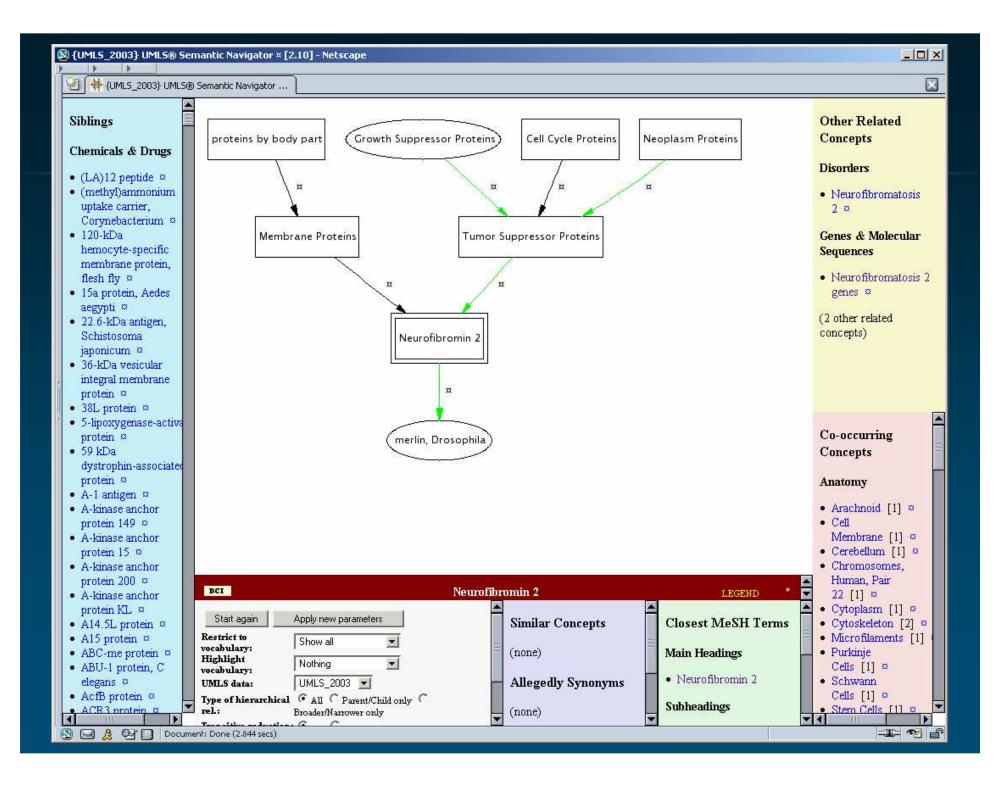


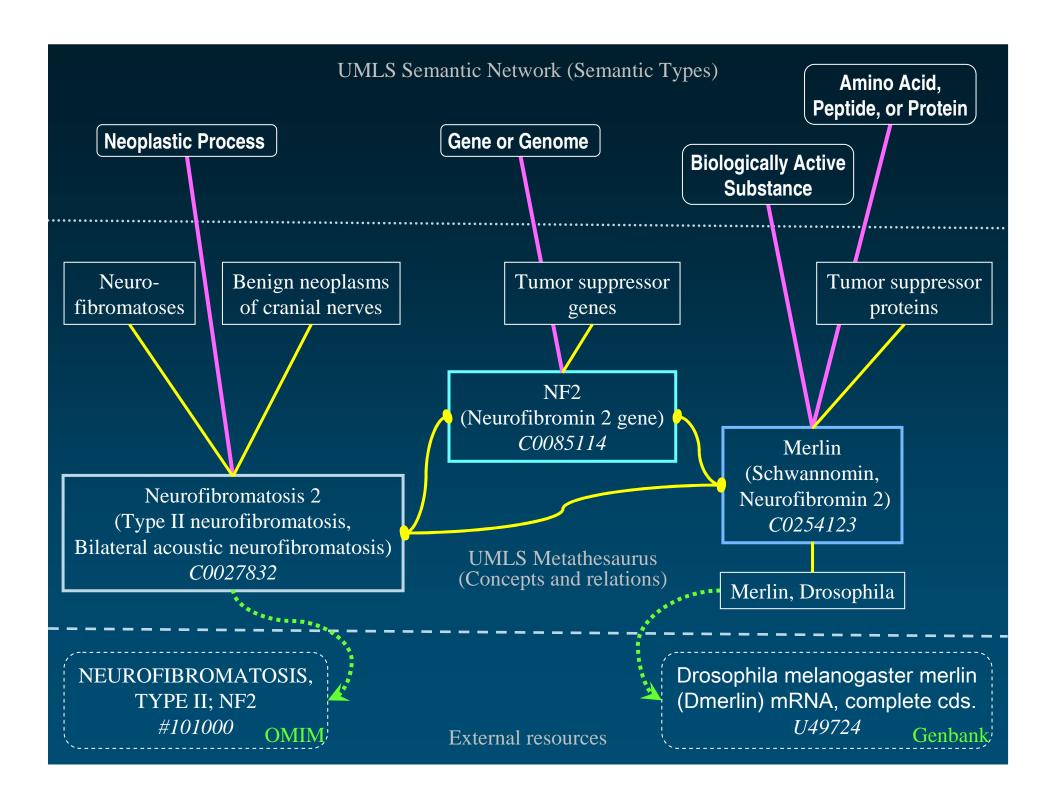


Merlin

- **♦** Synonyms
 - Neurofibromin 2
 - Schwannomin
 - Schwannomerlin
 - Neurofibromatosis-2
- ◆ 10 isoforms
- **♦** Annotations
 - Negative regulation of cell proliferation
 - Cytoskeleton
 - Plasma membrane







Limitations

- ◆ Genes not systematically represented
 - Most gene products and diseases are
- ◆ Gene/Gene product-Disease relations
 - Not systematically represented
 - Not explicitly represented (e.g., co-occurrence)
- Cross-references not systematically represented
- ◆ Naming conventions (genes)



Applications (1)

 $GenesTrace^{TM}$

Lussier Lab Columbia University

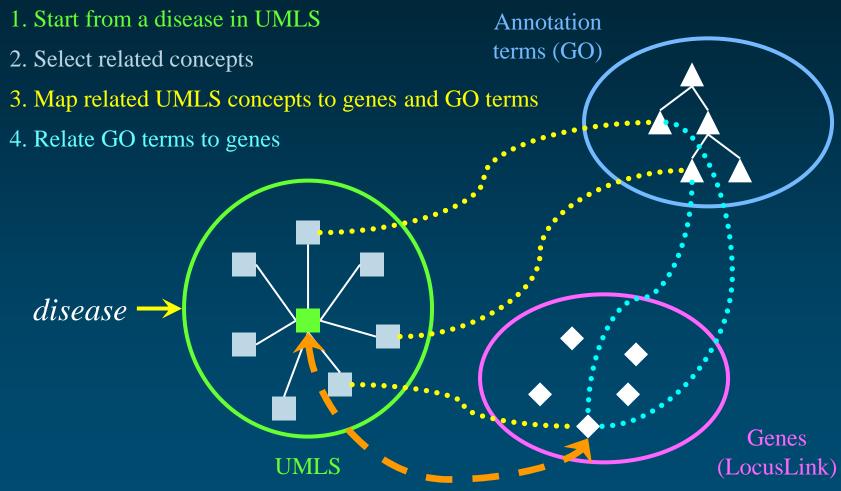
Objectives

◆ Relate diseases to genes through structured, integrated terminologies

◆ Biological Knowledge Discovery

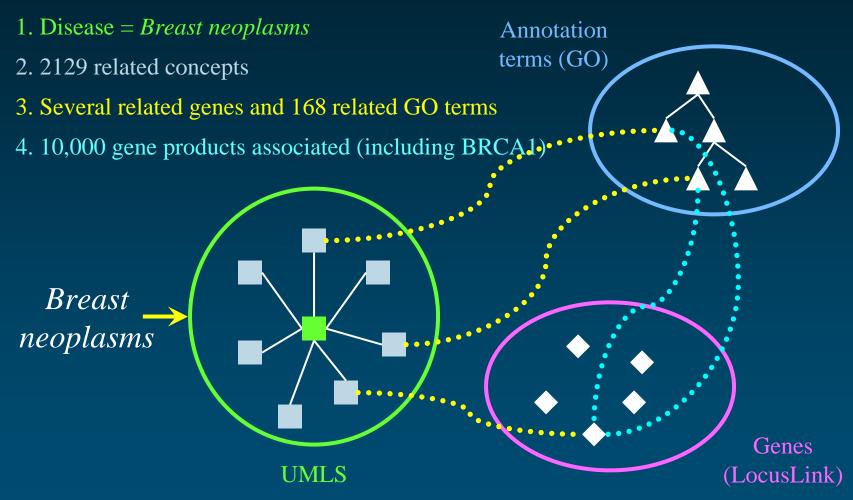


Resources and Methods





Validation Breast cancer – BRCA1 association





Limitations

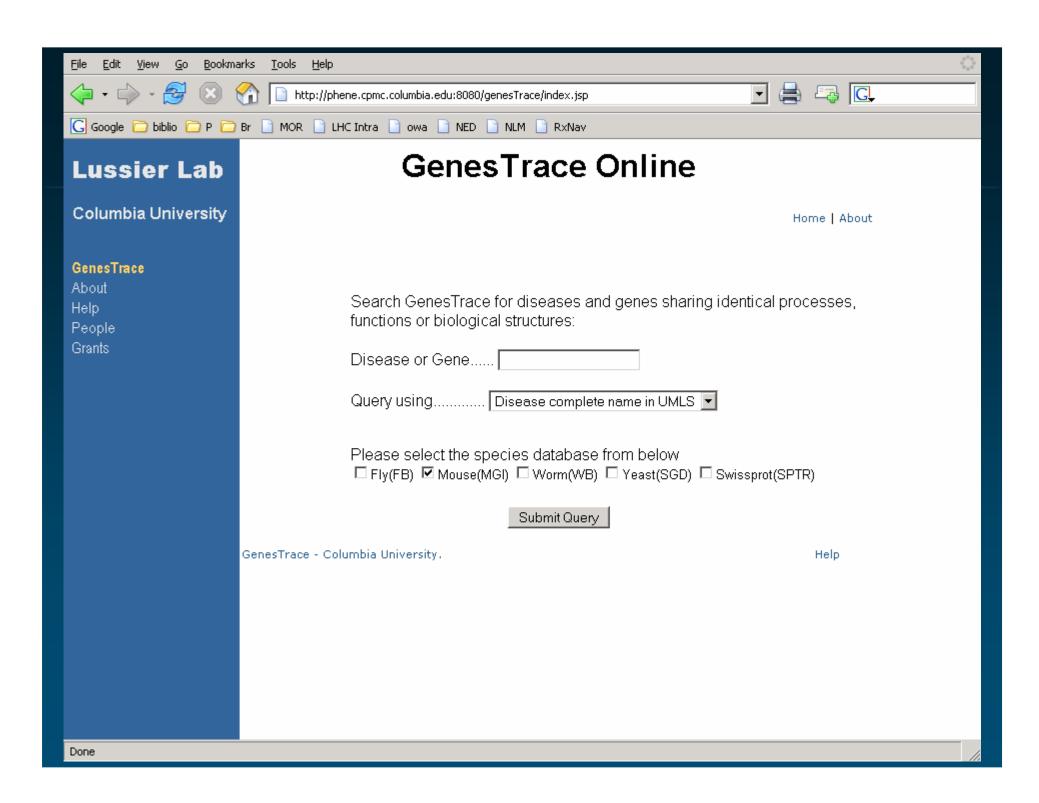
♦ Noise

- Too many non-specific GO terms associated (e.g., *nucleus*)
- Too many genes associated

♦ But

- Promising preliminary results
- Room for refinement





Applications (2)

$\overline{BioMeKe}$

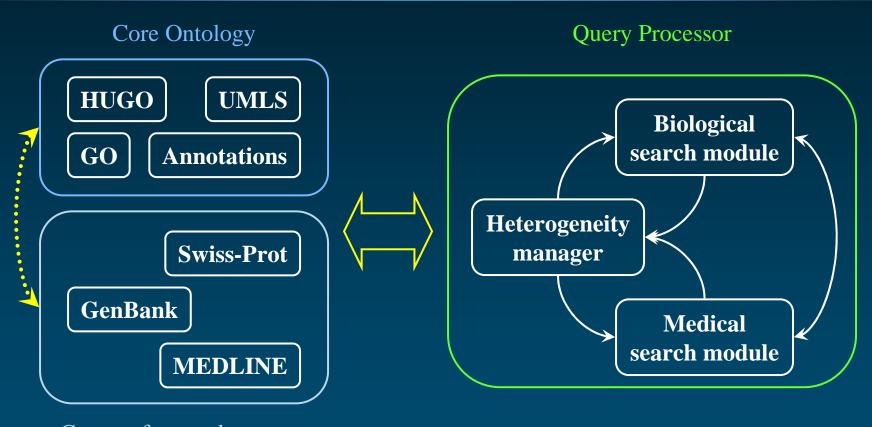
G. Marquet & al. LIM, Univ. Rennes, France

Objectives

- ◆ To develop a knowledge warehouse for transcriptome analysis (liver diseases)
- **◆** Semantic interoperability
 - Medical knowledge bases
 Clinical genomics
 - Molecular biology and genetics knowledge bases Functional genomics



Components





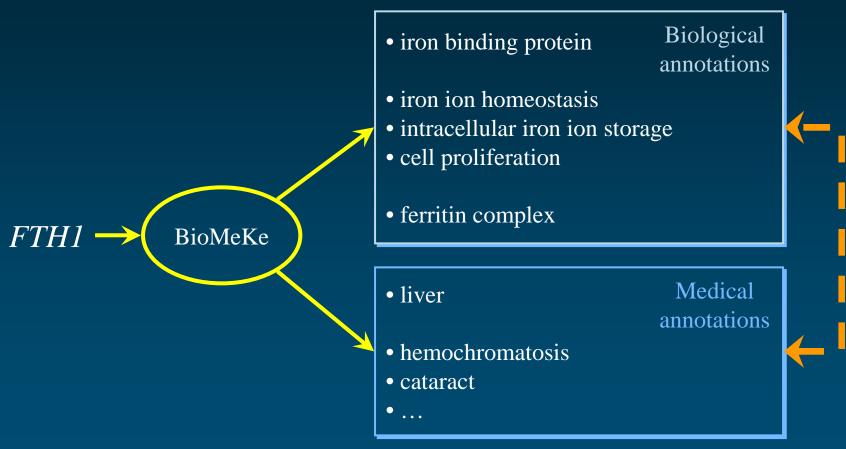


Example

- ◆ Input: ferritin, heavy polypedpide 1
- Mapping to biological resources
 - Not found in the Core ontology
 - Official name Ferritin heavy chain found through Xref
- ◆ Biological information obtained from GOA
- ◆ Mapping to medical resources
 - Not found in UMLS
 - Synonym *Ferritin H* found through Xref (Swiss-Prot)
- ◆ Medical information obtained through cooccurrence of MeSH index terms in MEDLINE



Results

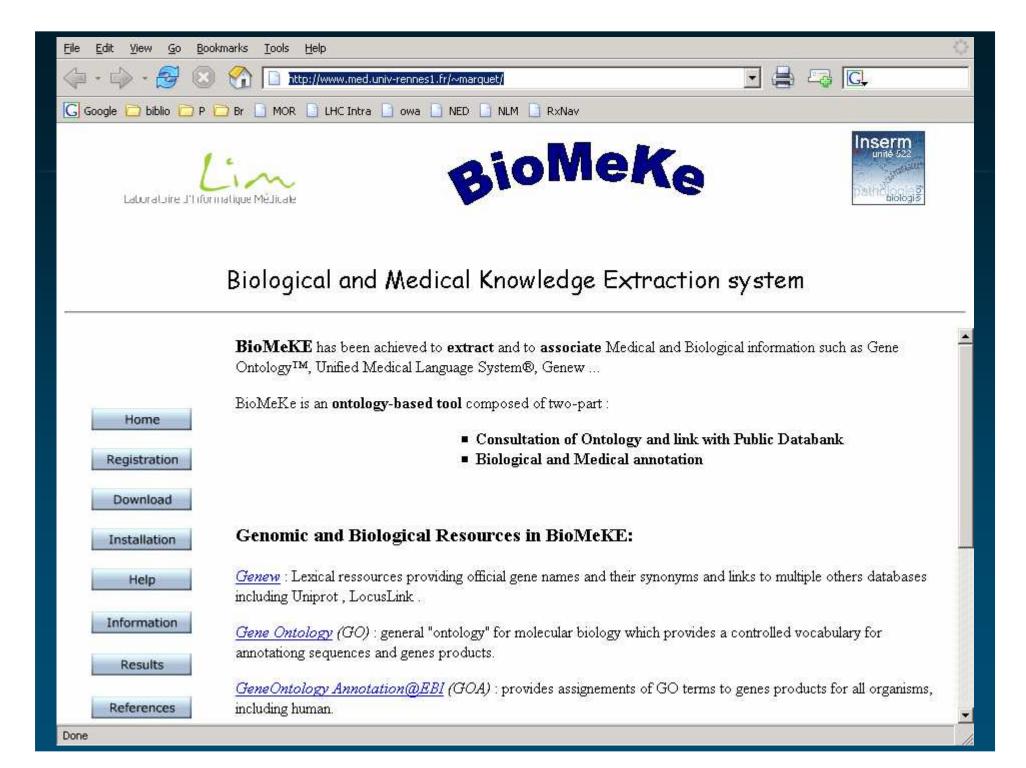




Limitations

- ◆ Non-formal ontologies
 - Knowledge may be inconsistently represented
 - Knowledge may be implicit (mappings)
- ◆ Partial automation
 - User input required to select databanks, reformulate queries
- **♦** Semantic integration
 - Naming issues
 - Mappings must be updated regularly



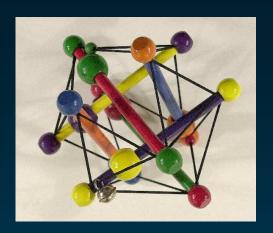


Conclusions

Conclusions

- ◆ Terminology integration provides some degree of information integration
- ◆ Most terminologies and the cross-referenced databases are readily available
- ◆ Lack of consistent representation
- Additional resources/techniques needed





Medical Ontology Research

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Questions

- ◆ What do I need to do to get the UMLS?
- ◆ What is an ontology?
- ◆ How is ontology different from
 - Terminology? / Database? / Knowledge base?
- ◆ Is the UMLS an ontology?
- ◆ Does the UMLS use Protégé?
- ◆ I heard of OWL. Is that any good?
- ◆ What is the Semantic Web going to do for us?



References UMLS

- ◆ UMLS umlsinfo.nlm.nih.gov
- **♦** UMLS browser
 - Knowledge Source Server: umlsks.nlm.nih.gov
 - Semantic Navigator: http://mor.nlm.nih.gov/perl/semnav.pl
 - (free, but UMLS license required)
- ◆ UMLS and information integration
 - O. Bodenreider. The UMLS: Integrating biomedical terminology. *Nucl. Acids Res.* 2004;32(1) (in press)



References Applications

♦ GenesTrace

- Cantor MN, Sarkar IN, Bodenreider O, Lussier YA. GenesTrace: Phenomic knowledge discovery via structured terminologies. In: Pacific Symposium on Biocomputing 2005; 2005. (in press).
- http://phene.cpmc.columbia.edu:8080/ genesTrace/index.jsp

♦ BioMeKE

- Marquet G, Burgun A, Moussouni F, Guerin E, Le Duff F, Loreal O. BioMeKe: an ontology-based biomedical knowledge extraction system devoted to transcriptome analysis. Stud Health Technol Inform. 2003;95:80-5.
- http://www.med.univ-rennes1.fr/~marquet/

